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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/462,909D

DATE: 08/09/2002
TIME: 13:51:05

Input Set : A:\65691179.app
Output Set: N:\CRF4\08092002\I462909D.raw

3 <110> APPLICANT: MEINIEL, ANNIE
 4 MONNERIE, HUBERT
 5 GOBRON, STEPHANIE
 7 <120> TITLE OF INVENTION: NOVEL POLYPEPTIDES AND POLYPEPTIDES USEFUL FOR
 8 REGENERATING THE NERVOUS SYSTEM
 10 <130> FILE REFERENCE: 065691/0179
 12 <140> CURRENT APPLICATION NUMBER: 09/462,909D
 13 <141> CURRENT FILING DATE: 2000-02-14
 15 <150> PRIOR APPLICATION NUMBER: PCT/FR98/01556
 16 <151> PRIOR FILING DATE: 1998-07-16
 18 <150> PRIOR APPLICATION NUMBER: FR 97/09016
 19 <151> PRIOR FILING DATE: 1997-07-16
 21 <160> NUMBER OF SEQ ID NOS: 168
 23 <170> SOFTWARE: PatentIn Ver. 2.1
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 37 <223> OTHER INFORMATION: Any amino acid; this range may encompass 1-5 amino
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194 <221> NAME/KEY: MOD_RES

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197 or no residues at all.
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205 <221> NAME/KEY: MOD_RES
206 <222> LOCATION: (9)
207 <223> OTHER INFORMATION: Not present if residue 8 is Pro; this residue is
208 Trp if residue 8 is Gly, Ser or Cys
210 <220> FEATURE:
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212 <222> LOCATION: (10)..(11)
213 <223> OTHER INFORMATION: Not present if residue 8 is Pro; these residues
214 are selected from Gly, Ser or Cys if residue 8 is
215 Gly, Ser or Cys
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220 <223> OTHER INFORMATION: Arg Ser, Val Ser or Val Thr
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223 <221> NAME/KEY: MOD_RES
224 <222> LOCATION: (18)..(22)
225 <223> OTHER INFORMATION: Any amino acid; this range may encompass 1-5 residues
226 or no residues at all.
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233 20
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299 Trp Gly Pro Cys Ser Val Ser Cys Gly
300 1 5
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314 1 5
317 <210> SEQ ID NO: 13
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320 <213> ORGANISM: Artificial Sequence
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

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Seq#:6; Xaa Pos. 1,4
Seq#:7; Xaa Pos. 3,6,9,10 // // // /
Seq#:9; Xaa Pos. 1,2,3,4,5,8,9,10,11,14,15,18,19,20,21,22
Seq#:10; N Pos. 6,9,15,18,24,27,30,36
Seq#:22; Xaa Pos. 2,3
Seq#:23; Xaa Pos. 1,2,3,4
Seq#:24; Xaa Pos. 3
Seq#:25; Xaa Pos. 3,6
Seq#:26; Xaa Pos. 3,6,7
Seq#:27; Xaa Pos. 3,6,7,8
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Seq#:30; Xaa Pos. 3,4,7
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